

OIPE

DATE: 04/08/2002 RAW SEQUENCE LISTING TIME: 15:11:42 PATENT APPLICATION: US/10/010,050A

Input Set : A:\97-38ClSequence Listing.txt Output Set: N:\CRF3\04082002\J010050A.raw

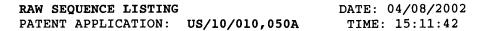


```
4 <110> APPLICANT: Sheppard, Paul O.
              Gilbertson, Debra G.
      7 <120> TITLE OF INVENTION: SECRETED PROTEINS ENCODED BY HUMAN
              CHROMOSOME 13
     11 <130> FILE REFERENCE: 97-38C1
                                                                   ENTERED
     13 <140> CURRENT APPLICATION NUMBER: US 10/010,050A
C--> 14 <141> CURRENT FILING DATE: 2002-03-26
     16 <150> PRIOR APPLICATION NUMBER: US 09/122,383
     17 <151> PRIOR FILING DATE: 1998-07-24
     19 <150> PRIOR APPLICATION NUMBER: US 60/053,613
     20 <151> PRIOR FILING DATE: 1997-07-24
     22 <160> NUMBER OF SEQ ID NOS: 19
     24 <170> SOFTWARE: FastSEQ for Windows Version 4.0
     26 <210> SEQ ID NO: 1
     27 <211> LENGTH: 1486
     28 <212> TYPE: DNA
     29 <213> ORGANISM: Homo sapien
     31 <220> FEATURE:
     32 <221> NAME/KEY: CDS
     33 <222> LOCATION: (47)...(1084)
     35 <400> SEQUENCE: 1
                                                                               55
     36 gaattcggca cgagggcagg aggtagacac ggcacagggc gccgag atg cgg cgg
                                                           Met Arg Arg
     37
     38
                                                                              103
     40 ggc gcg ggc gcg gct cgg gga cgc gct tcc tgg tgc tgg gcg ctg gcg
     41 Gly Ala Gly Ala Ala Arg Gly Arg Ala Ser Trp Cys Trp Ala Leu Ala
                                 10
             5
                                                                              151
     44 ctg ctt tgg ctc gcg gtg gtt ccg ggc tgg tcc cgg gtc tcg ggc atc
     45 Leu Leu Trp Leu Ala Val Val Pro Gly Trp Ser Arg Val Ser Gly Ile
                                                 30
     48 ccc tcc cgg cgc cac tgg ccg gtg ccc tac aag cgc ttt gac ttc cgt
                                                                              199
     49 Pro Ser Arg Arg His Trp Pro Val Pro Tyr Lys Arg Phe Asp Phe Arg
                                             45
     50
                         40
                                                                              247
     52 cca aaa cct gat cct tat tgt caa gct aag tat act ttc tgt cca act
     53 Pro Lys Pro Asp Pro Tyr Cys Gln Ala Lys Tyr Thr Phe Cys Pro Thr
                     55
                                         60
     56 ggc tca cct atc cca gtt atg gag ggt gat gat gac att gaa gtt ttt
                                                                              295
     57 Gly Ser Pro Ile Pro Val Met Glu Gly Asp Asp Ile Glu Val Phe
                                     75
                 70
                                                                              343
     60 cga tta caa gcc cca gta tgg gaa ttt aaa tat gga gac ctc ctg gga
     61 Arg Leu Gln Ala Pro Val Trp Glu Phe Lys Tyr Gly Asp Leu Leu Gly
                                 90
     62
             85
     64 cac ttq aaa att atg cat gat gcc att gga ttc aga agt aca tta act
                                                                              391
```

RAW SEQUENCE LISTING DATE: 04/08/2002 PATENT APPLICATION: US/10/010,050A TIME: 15:11:42

Input Set: A:\97-38ClSequence Listing.txt
Output Set: N:\CRF3\04082002\J010050A.raw

		Leu	Lys	Ile	Met	_	Asp	Ala	Ile	Gly		Arg	Ser	Thr	Leu		
	100					105					110					115	
				tac													439
69	Gly	Lys	Asn	${ t Tyr}$	Thr	Met	Glu	\mathtt{Trp}	Tyr	Glu	Leu	Phe	Gln	Leu	Gly	Asn	
70					120					125					130		
72	tat	aca	ttt	CCC	cat	ctc	cqa	cct	gaa	atq	gat	qcc	cct	ttc	taa	tqt	487
				Pro													
74	0,10			135			9		140		E			145		-1-	
	22+	722	aaa	gct	aaa	tac	+++	+++		ana	att	αat	rat		CaC	taa	535
																	333
	ASII	GIII	_	Ala	АТА	Cys	Pile		GIU	GIY	TTE	ASP	_	vaı	птэ	111	
78			150					155					160				500
				ggg													583
81	Lys	Glu	Asn	Gly	Thr	Leu	Val	Gln	Val	Ala	Thr	Ile	Ser	Gly	Asn	Met	
82		165					170					175					
84	ttc	aac	caa	atg	gca	aag	tgg	gtg	aaa	cag	gac	aat	gaa	aca	gga	att	631
85	Phe	Asn	Gln	Met	Ala	Lys	Trp	Val	Lys	Gln	Asp	Asn	Glu	Thr	Gly	Ile	
86	180					185	_		_		190				_	195	
		tat	σασ	aca	taa	aat	gta	aaa	acc	agc	cca	gaa	aaq	aaa	qca	gag	679
				Thr													
90	-1-	-1-	014		200	•••		-,,		205			-1-	V-1	210		
	202	+ ~~	+++	gat		+=0	~ ~ ~	+ ~+	tac		+++	ata	tta.	agg		+++	727
				-			-										, 2,
	THE	тр	PHE	Asp	Ser	TYL	ASP	Cys		гуз	FIIE	Val	Leu		TIIL	rne	
94				215					220					225			775
		-	_	gct	-			-			_			-			775
	Asn	Lys		Ala	GLu	Pne	GIY		GIU	Pne	гĀг	Asn		GIU	Thr	Asn	
98			230					235					240				
			_					_		_						aat	823
101	. Туі		_	g Ile	Phe	Leu			: Gly	7 Glu	ı Pro			Leu	ı GlZ	/ Asn	
102		245					250					255					
																gcc	871
105	Glu	ı Thi	: Sei	r Val	. Phe	: Gly	Pro	Thr	: Gly	Asr	ı Lys	Thi	Leu	ı Gly	, Let	ı Ala	
106	260)				265	•				270)				275	
108	ata	aaa	a aga	a ttt	. tat	tac	ccc	ttc	aaa	ı cca	cat	: ttg	, cca	act	aaa	ı gaa	919
109) Ile	Lys	Arg	y Phe	. Tyr	Tyr	Pro	Phe	Lys	Pro	His	Leu	ı Pro	Thi	Lys	s Glu	
110)				280					285	;				290)	
112	ttt	cto	ı tto	a aqt	cto	ttq	caa	att	ttt	gat	gea	qto	att	gto	cac	aaa	967
																Lys	
114				295					300					305			
		. ++	tat			tat	aat	+++			t a a	r +++	· tta			, aaa	1015
	-			_					_						_	Lys	1010
118		1 1110	31(FILE	. 1 <u>7</u> 1	. ADI	315		гтул		, 1116	320		Met	- шуэ	
				-													1062
																aga	1063
				5 TTE	: гуз	TTE		_	GIU	i Git	r TTE			Pro) ITE	e Arg	
122		325		_			330					335					
				a cto					laaca	icct	taat	tcta	ict c	jetet	tttt	it	1114
			Thi	r Leu	Ser	_		<u>l</u>									
	340					345											
																cttagcc	1174
129	ttt	ctto	cctt	ggtg	rcata	aa g	ttaa	aatg	c ac	atca	igcag	, aat	tgct	gca	tatt	aacatc	1234



Input Set : A:\97-38ClSequence Listing.txt
Output Set: N:\CRF3\04082002\J010050A.raw

131 132 133 134	cagi aaaa ggco	ggtga ttate aatae	atc t gta q cct t tcg a	ttggi ggaco tcaaa ag	tttca ctttq aaata	aa ti	tcc	gagco gggto	c tt	tgtt: acag:	aata atag	tgga ata	agaai tggt	tta gtg	tggtt ccca	agcgag ccatat gatttt aatagc	1294 1354 1414 1474 1486
	<210																•
	<213				46												
	<212																
	<213					o sap	pien										
	<400		-														
		Arg	Arg	Gly	Ala	Gly	Ala	Ala	Arg	_	Arg	Ala	Ser	Trp	Cys	\mathtt{Trp}	
143			_		5			_	_	10					15		
	Ala	Leu	Ala		Leu	Trp	Leu	Ala		Val	Pro	Gly	Trp		Arg	Val	
145	_		_,	20	_	_	_		25	_		_	_	30	_	_,	
	ser	GTA		Pro	Ser	Arg	Arg		Trp	Pro	Val	Pro	_	Lys	Arg	Phe	
147	•	51	35				•	40		~	a 1.		45	_	-1	_,	
	Asp		Arg	Pro	Lys	Pro	_	Pro	Tyr	Cys	GIn		Lys	Tyr	Thr	Phe	
149	_	50	_,	~ 3	_	_	55	_				60	_	_	_		
		Pro	Thr	GIA	ser		тте	Pro	vaı	мет		GTĀ	Asp	Asp	Asp		
151		**- 1	Dh.		.	70		D	**- 1		75	51	-		a 3	80	
	GIU	val	Pne	Arg		GIn	Ата	Pro	vaı	_	GIU	Pne	гÃг	Tyr	Gly	Asp	
153	T	T	a 1	77.2 -	85	T	7 1.	37- 4	77.4 -	90	11-	71 -	01	Dh.	95	G	
155	ьeu	ьеu	GIY	100	Leu	гаг	тте	met		Asp	Ата	тте	GIY		Arg	ser	
	mb	T 011	mb m		T	7.50	M	mb	105	01.	m	Ш	~1	110	Dha	71 m	
	THE	ьеu	115	GIĀ	глх	ASII	TAL	120	met	GIU	Trp	туг		Leu	Phe	GIN	
157	T 011	C1**		Crro	mh∽	Dho	Dwo		T 011	7 ~~	Dwo	C1	125 Wat	N an	Ala	Dwo	
159	ьеu	130	ASII	Cys	THI	Pile	135	пір	ьeu	AIG	PIO	140	мес	ASP	Ата	PIO	
_	Dho		Cve	Δan	Gln	Glv		Δla	Cve	Dho	Dho		G1 v	Tla	Asp	λen	
	145	111	Cys	ASII	GIII	150	ALU	AIU	Cys	FIIC	155	GIU	GLY	116	пор	160	
		His	Trn	Lvs	Glu		Glv	Thr	T.e.11	Va l		Va 1	Δla	Thr	Ile		
163	,				165		011			170	0111	, 41			175	001	
	Glv	Asn	Met	Phe		Gln	Met	Ala	Lvs		Val	Lvs	Gln	Asp	Asn	Glu	
165	0-1			180		0			185			-12		190		014	
	Thr	Glv	Ile		Tvr	Glu	Thr	Trp		Val	Lvs	Ala	Ser		Glu	Lvs	
167		_	195	-	-			200			-		205			•	
168	Gly	Ala	Glu	Thr	Trp	Phe	Asp	Ser	Tyr	Asp	Cys	Ser	Lys	Phe	Val	Leu	
169	-	210			-		215		-	_	-	220	-				
170	Arg	Thr	Phe	Asn	Lys	Leu	Ala	Glu	Phe	Gly	Ala	Glu	Phe	Lys	Asn	Ile	
															Thr		
173				•	245	_				250		•			255	•	
174	Leu	Gly	Asn	Glu	Thr	Ser	Val	Phe	Gly	Pro	Thr	Gly	Asn	Lys	Thr	Leu	
175		_		260					265			_		270			
	Gly	Leu	Ala	Ile	Lys	Arg	Phe	Tyr	Tyr	Pro	Phe	Lys	Pro	His	Leu	Pro	
177	_		275		_	-		280	7			-	285				
178	Thr	Lys	Glu	Phe	Leu	Leu	Ser	Leu	Leu	Gln	Ile	Phe	Asp	Ala	Val	Ile	
179		290					295					300					
180	Val	His	Lys	Gln	Phe	Tyr	Leu	Phe	Tyr	Asn	Phe	Glu	Tyr	${\tt Trp}$	Phe	Leu	

RAW SEQUENCE LISTING DATE: 04/08/2002 PATENT APPLICATION: US/10/010,050A TIME: 15:11:42

Input Set : A:\97-38ClSequence Listing.txt
Output Set: N:\CRF3\04082002\J010050A.raw

	305 310 315 320	
	Pro Met Lys Phe Pro Phe Ile Lys Ile Thr Tyr Glu Glu Ile Pro Leu	
183	325 330 335	
	Pro Ile Arg Asn Lys Thr Leu Ser Gly Leu	
185		
	<210> SEQ ID NO: 3	
	<211> LENGTH: 18	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial Sequence	
	<220> FEATURE:	
	<223> OTHER INFORMATION: Oligonucleotide ZC976	
	<400> SEQUENCE: 3	
	cgttgtaaaa cgacggcc	18
	<210> SEQ ID NO: 4	
	<211> LENGTH: 17	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial Sequence	
	<220> FEATURE:	
	<223> OTHER INFORMATION: Oligonucleotide ZC447	
	<400> SEQUENCE: 4	
	taacaatttc acacagg	17
	<210> SEQ ID NO: 5	
	<211> LENGTH: 20	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial Sequence	
	<220> FEATURE:	
	<223> OTHER INFORMATION: Oligonucleotide ZC14487	
	<400> SEQUENCE: 5	~ ~
	gacttccgtc caaaacctga	20
	<210> SEQ ID NO: 6	
	<211> LENGTH: 20	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial Sequence	
	<220> FEATURE:	
	<223> OTHER INFORMATION: Oligonucleotide ZC14716	
	<400> SEQUENCE: 6	20
	aggggcatcc atttcaggtg	20
	<210> SEQ ID NO: 7	
	<211> LENGTH: 20	
	<212> TYPE: DNA	
	<pre><213> ORGANISM: Artificial Sequence</pre>	
	<pre><220> FEATURE:</pre>	
	<pre><223> OTHER INFORMATION: Oligonucleotide ZC14712</pre>	
	<400> SEQUENCE: 7	20
	atggctaaac caagagtctt	20
	<210> SEQ ID NO: 8	
	<211> LENGTH: 20	
	<212> TYPE: DNA	
240	<213> ORGANISM: Artificial Sequence	

RAW SEQUENCE LISTING DATE: 04/08/2002 PATENT APPLICATION: US/10/010,050A TIME: 15:11:42

Input Set: A:\97-38ClSequence Listing.txt
Output Set: N:\CRF3\04082002\J010050A.raw

	248	<220> FEATURE:	
		<223> OTHER INFORMATION: Oligonucleotide ZC14710	
		<400> SEQUENCE: 8	
		gggtgaaaca ggacaatgaa '	20
		<210> SEQ ID NO: 9	
		<211> LENGTH: 20	
		<212> TYPE: DNA	
		<213> ORGANISM: Artificial Sequence	
		<220> FEATURE:	
	260	<223> OTHER INFORMATION: Oligonucleotide ZC14488	
		<400> SEQUENCE: 9	
	263	ttatgcacca aggaagaaag	20
		<210> SEQ ID NO: 10	
	266	<211> LENGTH: 20	
	267	<212> TYPE: DNA	
	268	<213> ORGANISM: Artificial Sequence	
	270	<220> FEATURE:	
	271	<223> OTHER INFORMATION: Oligonucleotide ZC14711	
		<400> SEQUENCE: 10	
	274	ttttctccaa tcaccagcat	20
	276	<210> SEQ ID NO: 11	
	277	<211> LENGTH: 18	
	278	<212> TYPE: DNA	
	279	<213> ORGANISM: Artificial Sequence	
	281	<220> FEATURE:	
	282	<223> OTHER INFORMATION: Oligonucleotide ZC14430	
	284	<400> SEQUENCE: 11	
	285	gtacatttcc ccatctcc	18
	287	<210> SEQ ID NO: 12	
	288	<211> LENGTH: 18	
	289	<212> TYPE: DNA	
	290	<213> ORGANISM: Artificial Sequence	
		<220> FEATURE:	
		<223> OTHER INFORMATION: Oligonucleotide ZC14431	
		<400> SEQUENCE: 12	
		ccattttcct tccagtga	18
		<210> SEQ ID NO: 13	
		<211> LENGTH: 1038	
		<212> TYPE: DNA	
		<213> ORGANISM: Artificial Sequence	
		<220> FEATURE:	_
		<223> OTHER INFORMATION: Degenerate nucleotide sequence encoding zsig46)
	305	polypeptide of SEQ ID NO:2	
W>		<pre><221> NAME/KEY: misc_feature</pre>	
		<222> LOCATION: (1)(1038)	
		<223> OTHER INFORMATION: $n = A,T,C$ or G	
		<400> 13 > 13	60
		atgmgnmgng gngenggnge ngenmgnggn mgngenwsnt ggtgytggge nytngenytn ytntggytng engtngtnee nggntggwsn mgngtnwsng gnatheenws nmgnmgneay	60
W>	313	1 chessis endendence uddiredaman mandenmand angenecuma imidimiducal	L20

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 04/08/2002 PATENT APPLICATION: US/10/010,050A TIME: 15:11:43

Input Set: A:\97-38ClSequence Listing.txt
Output Set: N:\CRF3\04082002\J010050A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the $\langle 220 \rangle$ to $\langle 223 \rangle$ fields of each sequence which presents at least one n or Xaa.

```
Seq#:13; N Pos. 6,9,12,15,18,21,24,27,30,33,36,39,51,54,57,60,63,69,72,75
Seq#:13; N Pos. 78,81,84,90,93,96,99,102,108,111,114,117,126,129,132,141
Seq#:13; N Pos. 153,156,162,168,180,189,198,201,204,207,210,216,219,228,246
Seq#:13; N Pos. 252,255,261,264,267,285,291,294,297,303,321,327,333,336,339
Seq#:13; N Pos. 342,345,348,360,378,387,390,399,405,411,414,417,429,432,450
Seq#:13; N Pos. 453,456,471,483,501,504,507,510,516,519,522,528,531,552,561
Seq#:13; N Pos. 579,582,597,606,612,615,618,627,630,636,648,660,669,672,675
Seq#:13; N Pos. 678,690,693,702,705,726,735,738,747,753,756,762,765,771,774
Seq#:13; N Pos. 783,786,789,795,798,801,804,813,816,819,822,825,834,846,855
Seq#:13; N Pos. 990,1005,1008,1011,1017,1026,1029,1032,1035,1038
```